## **WEST Search History**

Hide Items	Restore	Clear	Cancel
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DATE: Wednesday, September 14, 2005

Hide?	Set Name	Query	<u>Hit</u> Count
	DB=P	GPB,USPT,USOC,EPAB,JPAB,DWPI; PLUR=YES; OP=OR	
	L21	L20 same (multiple or plurality) and sequential	26
Ċ	L20	(ligase or ligation) same (construct or construction or make or making or synthesis) same (surface or solid or support) same (gene or polynucleotide or oligonucleotide or DNA or RNA)	1223
	L19	(ligase or ligation) near (construct or construction or make or making or synthesis) near (surface or solid or support)	0
	L18	(ligase or ligation) same (construct or construction or make or making or synthesis) same (surface or solid or support)	1935
Ū	L17	template same (ligase or ligation) same (construct or construction or make or making) same (surface or solid or support)	53
	L16	sequential same (hybridization or hybridize) same (ligase or ligation) same (construct or construction or make or making)	30
	L15	(sequential near (hybridization or hybridize) same (ligase or ligation)) same (construct or construction or make or making)	0
	L14	sequential near (hybridization or hybridize or ligate or ligation) same (construct or construction or make or making)	41
	L13	sequential near (hybridization or hybridize or ligate or ligation)	344
	L12	L9 same template	34
	L11	L9 same short	2
	L10	L9 same sequential	0
	L9	(assemble or assembly or synthesize or synthesis or construction) near (gene or oligonucleotide or polynucleotide) near (ligase or enzyme or enzymatic or ligation)	456
	L8	(assemble or assembly or synthesize or synthesis or construction) same (gene or oligonucleotide or polynucleotide) same (ligase or enzyme or enzymatic or ligation)	32220
	L7	6479262.pn.	2
	L6	ligase near "anchor"	1
	L5	ligase near "anchor probe"	1
	L4	L3 not L2	5
	L3	(coupled or couple or linked or link or crosslink or "cross link" or "cross-link" or tether or tethered or anchor or anchored or bound or bind or connect) near ligase	220
	L2	(coupled or couple or linked or link or crosslink or "cross link" or "cross-link" or tether or anchor or anchored or bound or bind or connect) near ligase	215

## **END OF SEARCH HISTORY**

FILE 'USPATFULL, MEDLINE, CAPLUS' ENTERED AT 1:	1:35:58 ON 14 SE	P 2005
L1 24 S TETHER? (4A) LIGASE		
L2 2 S TETHER? (W) LIGASE		
L3 1856 S (CROSS(W) LINK OR COUPL? OR ANCHOR	OR LINK? OR BO	UND OR BIND)
L4 58 S (CROSS(W)LINK OR COUPL? OR ANCHOR	OR LINK? OR BO	UND OR BIND)
L5 52 DUP REM L4 (6 DUPLICATES REMOVED)		
L6 1684 DUP REM L3 (172 DUPLICATES REMOVED)		
=> log h		
•	NCE FILE TO	TAL
	ENTRY SESS	ION
FULL ESTIMATED COST	97.79 98	.21
DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS) SI	ICE FILE TO	TAL
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SESSION WILL BE HELD FOR 60 MINUTES		

SESSION WILL BE HELD FOR 60 MINUTES
STN INTERNATIONAL SESSION SUSPENDED AT 11:50:37 ON 14 SEP 2005

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NUMBER KIND DATE -----

US 6221600 B1 20010424 US 1999-414847 19991008 PATENT INFORMATION: 19991008 (9) APPLICATION INFO.:

DOCUMENT TYPE: Utility FILE SEGMENT: Granted

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NUMBER OF DRAWINGS: 30 Drawing Figure(s); 16 Drawing Page(s)

LINE COUNT: 3179

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention relates to a method for the detection of gene expression and analysis of both known and unknown genes. The invention is a highly sensitive, rapid and cost-effective means of monitoring gene expression, as well as for the analysis and quantitation of changes in gene expression for a defined set of genes and in response to a wide variety of events. It is an important feature of the present invention that no single molecular species of cDNA gives rise to more than one fragment in the collection of products which are subsequently amplified and representative of each expressed gene. This achievement is facilitated by immobilizing the cDNA prior to digesting and then digesting with sequentially with two frequently cutting enzymes. Linker oligomers are ligated to each cut site following the respective digestion. Primers, complementary to the oligomer sequence with an additional 3' variable sequence are used to amplify the fragments. Using an array of fragments theoretically facilitates the amplification of all of the possible messages in a given sample.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 52 OF 52 CAPLUS COPYRIGHT 2005 ACS on STN

ACCESSION NUMBER: 1986:16901 CAPLUS

DOCUMENT NUMBER: 104:16901

DNA ligase activity in pea seedlings (Pisum sativum TITLE:

L.): development of a sensitive assay system and partial characterization of soluble and chromatin-

bound ligases

AUTHOR(S): Daniel, Paul P.; Bryant, John A.; Barker, David G. Dep. Plant Sci., Univ. Coll., Cardiff, CF1 1XL, UK CORPORATE SOURCE: SOURCE:

Biochemistry International (1985), 11(5), 645-52

CODEN: BIINDF; ISSN: 0158-5231

DOCUMENT TYPE: Journal LANGUAGE: English

An efficient extraction procedure and a rapid and quant. assay for plant DNA ligases are described. By using these extraction and assay procedures, it was

possible for the 1st time to detect and partially characterize 2 populations of DNA ligase, one chromatin-bound and one soluble, in pea seedlings. The 2 populations of the enzyme showed similar assay

requirements and pH optima and were both stimulated by Nonidet P40 and by

PEG 6000. The chromatin-bound ligase was stimulated

by spermidine, whereas the soluble ligase was inhibited by this compound

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